RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/623,914A							
Source:	1FW16,							
Date Processed by STIC:	3/22/07							

ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 03/22/2007
PATENT APPLICATION: US/10/623,914A TIME: 15:27:05

Input Set : A:\27708465.APP

Output Set: N:\CRF4\03222007\J623914A.raw

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3 <110> APPLICANT: HERING, THOMAS M.
         JOHNSTONE, BRIAN
 6 <120> TITLE OF INVENTION: PROBES FOR CHONDROGENESIS
 8 <130> FILE REFERENCE: 27708/04065
10 <140> CURRENT APPLICATION NUMBER: 10/623,914A
11 <141> CURRENT FILING DATE: 2003-07-21
                                                      see p.6
13 <150> PRIOR APPLICATION NUMBER: 60/211,384
14 <151> PRIOR FILING DATE: 2000-06-14
16 <160> NUMBER OF SEQ ID NOS: 7
18 <170> SOFTWARE: PatentIn Ver. 3.3
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 2666
22 <212> TYPE: DNA
23 <213> ORGANISM: Homo sapiens
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28 gtcactggct gtcatgtagt ttcctcaact actgcctcag ctctacaatc ccagagtaaa 180
29 getettetee aaatgaagag eeaggaagag gtagaggtgg eaggaattaa aetttgtaaa 240
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32 aggaacctag tttcagtggg tctttgcatt tctaaaccag atgtgatctc cttactggag 420
33 caagagaaag accettgggt gataaaagga gggatgaaca gaggcetgtg eccagaettg 480
34 gagtgtgtgt gggtgaccaa atcattatct ttaaaccagg atatttatga agaaaaatta 540
35 cccccggcaa tcataatgga aagacttaaa agctatgacc ttgaatgttc aacattaggg 600
36 aaaaactgga aatgtgaaga cttgtttgag agggagcttg taaaccagaa gacacatttt 660
37 aggcaagaga ccatcactca tatagatact cttattgaaa aaagagatca ctctaacaaa 720
38 tetgggacag titticatet gaatacatta tettatataa aacagatitt teecatggaa 780
39 gagagaatat ttaattttca tacagataag aaaagcttaa aaacacattc agttgtgaaa 840
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41 ttcagcaaaa tctcaaccct tactcttcac caaagaattc atacaggaga gaaaccctat 960
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43 atacadacag gagaaaaacc ttttgaatgt actgaatgtg ggaaagcctt cagccagaat 1080
44 gctcatcttg ttcaacacca gagagttcat actggagaga aaccttatca gtgtaagcag 1140
45 tgtaataaag cattcagcca gcttgcacac cttgctcaac atcagagggt ccacactgga 1200
46 gagaaacct atgaatgtat tgaatgtggg aaggetttta gtgattgete atceetaget 1260
47 catcatcgaa ggattcacac tgggaaaaga ccttatgaat gtattgactg tgggaaagct 1320
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51 ggctcatctc tgacagtaca tcagagaatt catacaggag agaaacctta tgaatgcaat 1560
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Input Set : A:\27708465.APP

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57 agagttcata ctggtgagaa gccttacgaa tgtattgaat gtgggaaggc ctttagtgat 1920
58 ggctcatatc ttgttcaaca tccgagactc cacagtggca aaagaccgta tgaatgtctt 1980
59 gaatgtggga aggcattcag gcagagggca tccttgattt gtcatcagag atgtcatact 2040
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61 actctqcatc aqaqaattca tacaqqaqaq aaaccttatq aqtqtaaqqa atqtagcaaa 2160
62 geetteagee aggttgeeea tettaeteta eataagagaa tteataetgg agaaaggeee 2220
63 tatgagtgta aagaatgtgg aaaagccttc aggcagagtg tacatcttgc tcatcatcag 2280
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65 ctatagatte aatetegtaa atgettetag catecatetg etttttteea geacatgtee 2400
66 catcatcata gtccaagacg caaccatctc atctggattt ctgcagtagc ataactgttg 2460
67 ccccttttgc ttctatcaac tacatgttta acactgtagg cagcctaacc ttttaaaaat 2520
68 aaaaatacat aatttatgtt attttcccat ttaaaacact tgatttgaaa aatatattaa 2580
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75 <212> TYPE: PRT
76 <213> ORGANISM: Homo sapiens
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88 Lys Lys Val Met Leu Glu Asn Tyr Arg Asn Leu Val Ser Val Gly Leu
91 Cys Ile Ser Lys Pro Asp Val Ile Ser Leu Leu Glu Gln Glu Lys Asp
92 65
                        70
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94 Pro Trp Val Ile Lys Gly Gly Met Asn Arg Gly Leu Cys Pro Asp Leu
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97 Glu Cys Val Trp Val Thr Lys Ser Leu Ser Leu Asn Gln Asp Ile Tyr
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98
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100 Glu Glu Lys Leu Pro Pro Ala Ile Ile Met Glu Arg Leu Lys Ser Tyr
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103 Asp Leu Glu Cys Ser Thr Leu Gly Lys Asn Trp Lys Cys Glu Asp Leu
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109 Ile Thr His Ile Asp Thr Leu Ile Glu Lys Arg Asp His Ser Asn Lys
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112 Ser Gly Thr Val Phe His Leu Asn Thr Leu Ser Tyr Ile Lys Gln Ile
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115 Phe Pro Met Glu Glu Arg Ile Phe Asn Phe His Thr Asp Lys Lys Ser
118 Leu Lys Thr His Ser Val Val Lys Lys His Lys Gln Asp Arg Gly Glu
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Input Set : A:\27708465.APP

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		Thr	Leu	Thr	Leu	His	Gln	Arg	Ile	His	Thr	Gly	Glu	Lys	Pro	Tyr
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128		0,70		260	0,10	0-1	-1-		265					270		
	Gln	His	Gln		Tle	His	Thr	Glv		Lvc	Pro	Phe	Glu		Thr	Glu
131	01		275	*** 9	110			280	014				285	0,0		
	Cvc	Glv		Δ] =	Dha	Ser	Gln		בומ	Hic	T.011	Val		Hic	Gln	Ara
134	_	290	цуз	AIG	1110	DCI	295	HOII	ΑIQ	1115	±Cu.	300	Q111		01	9
			Thr	Clv	Glu	Lys		Тит	Gln	Cve	Lare		Cvc	Δen	T.vc	Δla
	305	птэ	1111	Gry	Giu	310	FIU	1 y 1	GIII	Cys	315	GIII	Cys	NO.II	цуз	320
		Com	C1 ~	T 011	7 J -	His	T 011	71-	C15	пiс		7 ~~	7727	uia	Thr	
	Pile	ser	GIII	ьeu		птѕ	Leu	Ala	GIII	330	GIII	Arg	vai	птэ		GIY
140	~1	.	D		325	a	71 -	a 1	a		T		Dh.	C	335	C
	GIU	ьys	PIO	_	GIU	Cys	шe	GIU		GIY	пуѕ	Ald	Pne		Asp	Cys
143	.	0	.	340	***	TT -	3	3	345	TT -	m1	a 1	T	350	D	
		ser		Ата	HIS	His	Arg	_	тте	HIS	Thr	GIY	_	Arg	PIO	Tyr
146.		_	355		~	~ 3	_	360	-	_	~ 3	_	365	a	.	~ 7 -
	GIu	_	Пе	Asp	Cys	Gly	-	Ala	Pne	Arg	Gin		Ата	Ser	ьeu	TIE
149	_	370	_	_	_		375		~1	~ 3	_	380	-1	_	~	~7.
	-	HIS	Arg	Arg	ıyr	Tyr	HIS	Thr	GIA	GIU	_	Pro	Pne	Asp	Cys	
	385	_		_		390	1	_			395		-1	a 1 .		400
	Asp	Cys	GIY	Lys		Phe	Thr	Asp			GIY	ьеu	11e	Gin		гàг
155			•		405			_^		410	_	_		_	415	_
	Arg	Ile	His		GIY	Glu	Arg	Pro		Lys	Cys	Asn	Val		Gly	гàг
158				420		_		_	425				_	430	1	
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161			435					440				_	445			•
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173				500					505					510		
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176			515					520					525			
178	Lys	Ile	His	Thr	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Lys	Glu	Arg	Gly	Lys
179		530					535					540				
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184	Gly	Glu	Lys	Pro	Tyr	Glu	Cys	Ile	Glu	Cys	Gly	Lys	Ala	Phe	Ser	Asp
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Input Set : A:\27708465.APP

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199 Arg Ile His Thr Gly Glu Lys Pro Tyr Glu Cys Lys Glu Cys Ser Lys
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205 Gly Glu Arg Pro Tyr Glu Cys Lys Glu Cys Gly Lys Ala Phe Arg Gln
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223 gatgtaatgt tggagaacta tagtaacttg gtgtcactgg atttggagtc aaaaacgtat 180
224 gagaccaaaa aatatttttc agaaaatgat atttttgaaa taaatttttc ccagtgggag 240
225 atgaaggaca aaagtaaaac ccttggcctt gaggcatcca tcttcagaaa taattggaag 300
226 tgcaaaagca tattcgaggg actaaaagga catcaagagg gatacttcag tcaaatgata 360
227 atcagctatg aaaaaatacc ttcttacaga aaaagtaaat ctcttactcc acatcaaaga 420
228 attcataata cagagaaatc ctatgtttgt aaggaatgtg ggaaggcttg cagtcatggc 480
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233 tttagtcgtg gctatcacct tacccaacat cagaaaattc atattggtgt gaaatcttat 780
234 aaatgtaagg aatgtgggaa ggcctttttt tggggctcaa gccttgctaa acatgagata 840
235 attcatacag gtgagaaacc ttataaatgt aaagaatgtg ggaaggcctt cagtcgtggc 900
236 tatcaactta ctcaqcatca qaaaatccat actggtaaga aaccttatga atgtaaaata 960
237 tgtggaaagg ctttttgttg gggctatcaa cttactcgac atcagatatt tcatactggt 1020
238 gagaaaccct atgaatgcaa ggaatgtggg aaggctttta attgcggatc aagtcttatt 1080
239 caacatgaaa gaattcatac tggtgagaaa ccttatgaat gtaaagaatg tggaaaggcc 1140
240 tttagtcgtg gctatcacct ttctcaacat cagaaaatcc atactggtga gaaacctttt 1200
241 gaatgtaagg aatgtgggaa ggcctttagt tggggttcaa gccttgttaa acatgagaga 1260
242 gttcatactg gtgagaaatc ccatgaatgt aaagaatgcg gaaagacctt ttgtagtggg 1320
243 tatcaactta ctcgacatca ggtatttcac actggtgaga aaccctatga atgtaaggaa 1380
244 tgtgggaagg cttttaattg tggatcaagc cttgttcaac atgaaagaat ccatacaggg 1440
245 gagaaaccct atgaatgtaa agaatgtgga aggcttttag tcgtggctat caccttactc 1500
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248 aatgtaaaga ctgtgggaag gcctttggta gtggctatca acttagtgtt catcagagat 1680
249 ttcatactgg tgagaagctt tatcaacata aggaattcgg gaagaccttt actcgtggct 1740
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Input Set : A:\27708465.APP

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254 tatgggcaat tatcttgcta tccagcaatt catactagtg agaaatattt tgaatataat 2040
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271 Asp Val Met Leu Glu Asn Tyr Ser Asn Leu Val Ser Leu Asp Leu Glu
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                                 40
274 Ser Lys Thr Tyr Glu Thr Lys Lys Ile Phe Ser Glu Asn Asp Ile Phe
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277 Glu Ile Asn Phe Ser Gln Trp Glu Met Lys Asp Lys Ser Lys Thr Leu
278 65
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                                              75
280 Gly Leu Glu Ala Ser Ile Phe Arg Asn Asn Trp Lys Cys Lys Ser Ile
283 Phe Glu Gly Leu Lys Gly His Gln Glu Gly Tyr Phe Ser Gln Met Ile
284
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                                    105
286 Ile Ser Tyr Glu Lys Ile Pro Ser Tyr Arg Lys Ser Lys Ser Leu Thr
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289 Pro His Gln Arg Ile His Asn Thr Glu Lys Ser Tyr Val Cys Lys Glu
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292 Cys Gly Lys Ala Cys Ser His Gly Ser Lys Leu Val Gln His Glu Arg
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295 Thr His Thr Ala Glu Lys His Phe Glu Cys Lys Glu Cys Gly Lys Asn
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                                        170
298 Tyr Leu Ser Ala Tyr Gln Leu Asn Val His Gln Arg Phe His Thr Gly
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301 Glu Lys Pro Tyr Glu Cys Lys Glu Cys Gly Lys Thr Phe Ser Trp Gly
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304 Ser Ser Leu Val Lys His Glu Arg Ile His Thr Gly Glu Lys Pro Tyr
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307 Glu Cys Lys Glu Cys Gly Lys Ala Phe Ser Arg Gly Tyr His Leu Thr
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                                            235
310 Gln His Gln Lys Ile His Ile Gly Val Lys Ser Tyr Lys Cys Lys Glu
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313 Cys Gly Lys Ala Phe Phe Trp Gly Ser Ser Leu Ala Lys His Glu Ile
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316 Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Lys Glu Cys Gly Lys Ala
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/623,914A

DATE: 03/22/2007 TIME: 15:27:06

Input Set : A:\27708465.APP

Output Set: N:\CRF4\03222007\J623914A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 6,9,18

Seq#:7; Xaa Pos. 1,2,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23 Seq#:7; Xaa Pos. 24,25,26,27,29,30,31,33,34,35,36,37,39,40,42,43,44,45,46

VERIFICATION SUMMARY

DATE: 03/22/2007

PATENT APPLICATION: US/10/623,914A

TIME: 15:27:06

Input Set : A:\27708465.APP

Output Set: N:\CRF4\03222007\J623914A.raw

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